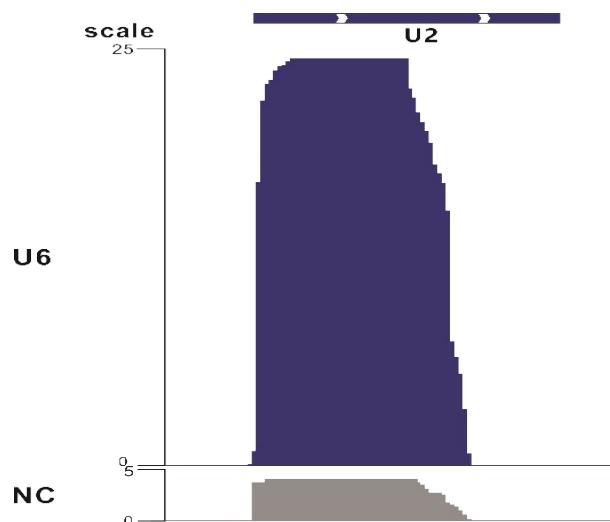


**Electronic Supplementary Information (ESI)**

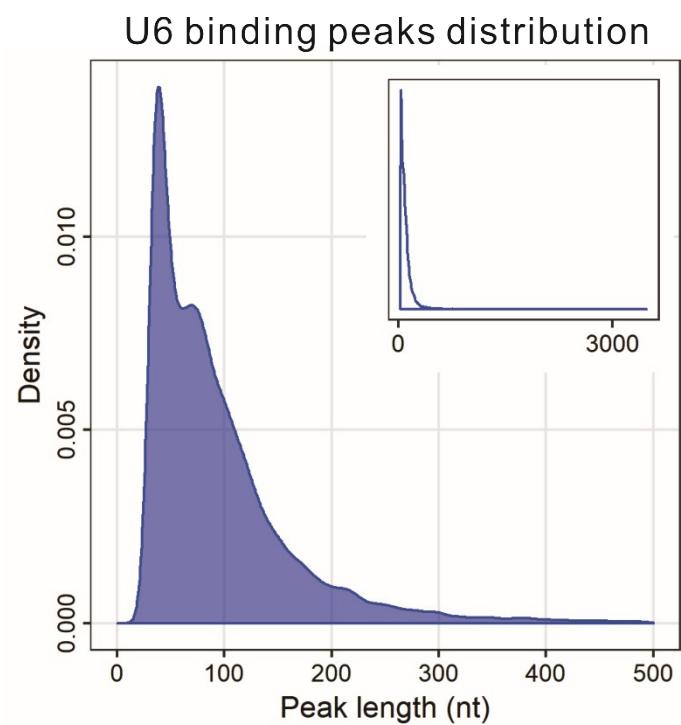
**CIRDES: an efficient genome-wide method for *in vivo* RNA-RNA interactome analysis**

Yao-Ting Li<sup>\*a</sup>, Nan Zhou<sup>\*a</sup>, Wei-Xi Deng<sup>\*a</sup>, Xue-Zhen Zeng<sup>b</sup>, Xiao-Juan Wang<sup>a</sup>, Jing-Wen Peng<sup>a</sup>, Bin Yang<sup>a</sup>, Yan-Jie Wang<sup>a</sup>, Jian-You Liao<sup>#a</sup>, Dong Yin<sup>#a</sup>

<sup>a</sup> Guangdong Provincial Key Laboratory of Malignant Tumor Epigenetics and Gene Regulation, Medical Research Center, Sun Yat-Sen Memorial Hospital, Sun Yat-Sen University, Guangzhou 510120, China. E-mail: [liaojy3@mail.sysu.edu.cn](mailto:liaojy3@mail.sysu.edu.cn), [yind3@mail.sysu.edu.cn](mailto:yind3@mail.sysu.edu.cn); Tel: +86-20-89205524  
<sup>b</sup> Department of Liver Surgery, The First Affiliated Hospital of Sun Yat-Sen University, Guangzhou 510120, China.



Supplementary Fig. 1 U6 binding sites on U2.



Supplementary Fig. 2 The length distribution of U6 binding sites.